

```
<80..241
    Region
                     /region name="LRR RI"
                     /note="Leucine-rich repeats (LRRs), ribonuclease inhibitor
                     (RI)-like subfamily. LRRs are 20-29 residue sequence
                     motifs present in many proteins that participate in
                     protein-protein interactions and have different functions
                     and cellular locations; cd00116"
                     /db xref="CDD:29015"
    Site
                     186
                     /site_type="glycosylation"
                     /experiment="experimental evidence, no additional details
                     /citation=[1]
    Region
                     299..346
                     /region_name="LRRCT"
                     /note="Leucine rich repeat C-terminal domain; smart00082"
                     /db xref="CDD:47408"
                     299..346
    Region
                     /region_name="Leucine rich repeat C-terminal domain"
                     /note="LRRCT; Leucine rich repeat C-terminal domain"
                     /db xref="CDD:smart00082"
                     299..346
                     /region name="Leucine rich repeat C-terminal domain.
                     Leucine Rich Repeats pfam00560 are short sequence motifs
                     present in a number of proteins with diverse functions and
                     cellular locations. Leucine Rich Repeats are often flanked
                     by cysteine rich domains. This domain is often found at
                     the C-terminus of tandem leucine rich repeats"
                     /note="LRRCT; Leucine rich repeat C-terminal domain.
                     Leucine Rich Repeats pfam00560 are short sequence motifs
                     present in a number of proteins with diverse functions and
                     cellular locations. Leucine Rich Repeats are often flanked
                     by cysteine rich domains. This domain is often found at
                     the C-terminus of tandem leucine rich repeats"
                     /db_xref="CDD:pfam01463"
                     1..347
    CDS
                     /gene="LRG"
                     /coded_by="NM_052972.1:12..1055"
                     /note="contains leucine-rich repeat"
                     /db_xref="GeneID:116844"
ORIGIN
        1 msswsrgrpk spggigphys rtlfllllla asawgytlsp kdcgyfrsdh gssiscgppa
       61 eipgylpadt vhlaveffnl thipanligg askigelhis snglesispe firpvpglrv
      121 ldltrnaltg lppglfgasa tldtlvlken glevlevswl hglkalghld lsgnrlrklp
      181 pgllanftll rtldlgengl etlppdllrg plqlerlhle gnklqvlgkd lllpqpdlry
      241 lflngnklar vaagafqqlr qldmldlsnn slasvpeqlw aslgqpnwdm rdgfdisgnp
      301 wicdqnlsdl yrwlgagkdk mfsqndtrca gpeavkggtl lavaksg
```

Disclaimer | Write to the Help Desk NCBI | NLM | NIH

lun 19 2007 (3.56:00